



Figure 2

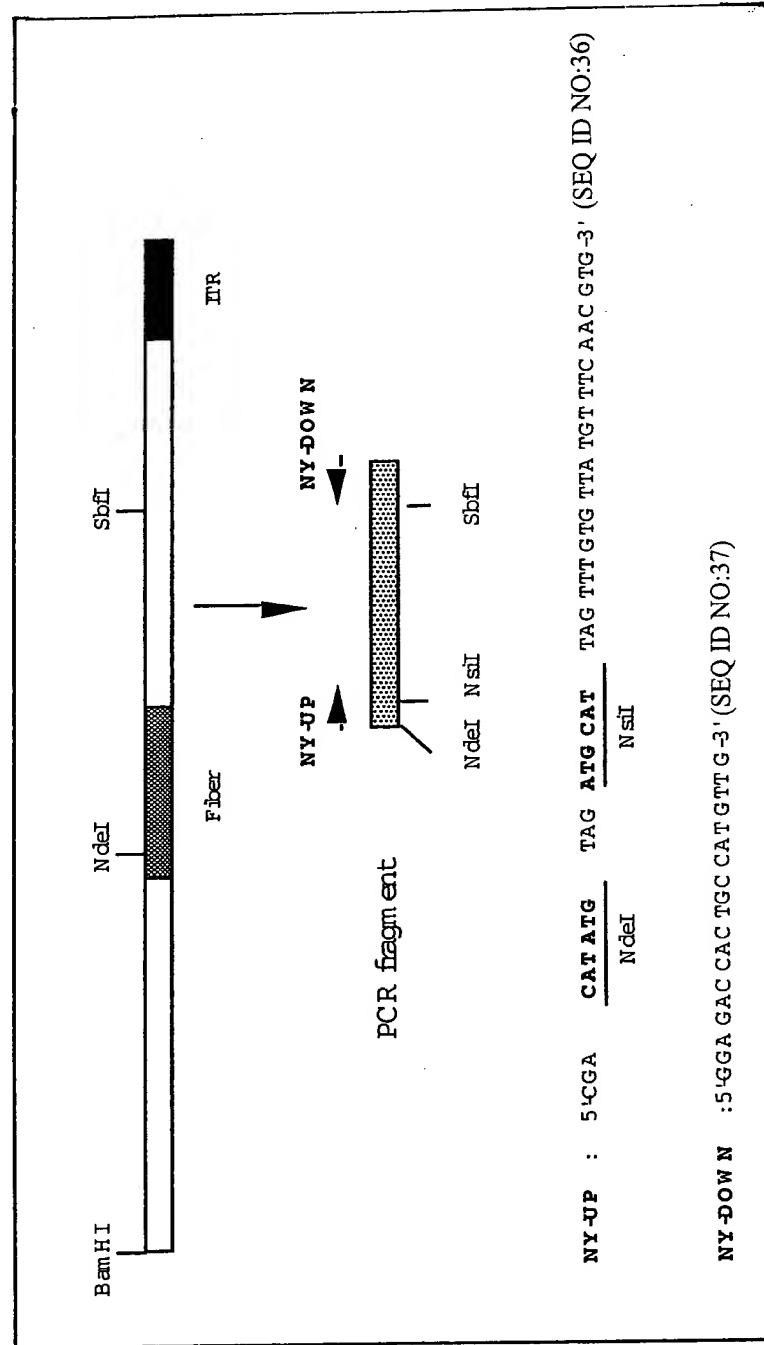




Figure 4: Sequence of Ad5/fib16 chimeric fiber

ATGAAGCGCGCAAGACCGTCTGAAGATACTTCACCCGTGTATCCATATGAAGATGAAAGCAGCT
CACAAACACCCCTTATAAACCTGGTTCATTCCTCAAATGGTTTGACAAAGCCCAGATGGAGT
TCTAACTCTTAAATGTGTTAATCCACTCACTACCGCCAGCGGACCCCTCCAACCTAAAGTTGGAAGC
AGTCTTACAGTAGATACTATCGATGGTCTTGGAGGAAAATAACTGCCGAAGCGCCACTCACTA
AAACTAACCACTCCATAGGTTATTAATAGGATCTGGCTTGCAACAAAGGATGATAAAACTTTGTTT
ATCGCTGGGAGATGGGTGTAACAAAGGATGATAAAACTATGTTATCGCTGGGAGATGGGTTAATA
ACAAAAAAATGATGTAATGTGCCAAACTAGGACATGGCCTGTGTTGACTCTTCAATGCTATCA
CCATAGAAAACAACACCTTGTGGACAGCGCAAAACCAAGCGCAACTGTGTAATTAAAGAGGGAGA
AGATTCCCCAGACTGTAAGCTCACTTAGTTAGTGAAGAATGGAGGACTGATAAAATGGATACATA
ACATTAATGGGAGCCTCAGAATATACTAACACCTTGTAAAACAAATCAAGTTACAATCGATGTA
ACCTCGCATTTGATAATACTGGCCAAATTATTACTTACCTATCCCTTAAAGTAACCTGAACCT
TAAAGACAACCAAAACATGGCTACTGGAACCATACCAGTGCCAAAGGCTTCATGCCAGCACC
GCCTATCCATTATAACATACGCCACTGAGACCCCTAAATGAAGATTACATTATGGAGAGTGTACT
ACAAATCTACCAATGGAACCTCTTCCACTAAAGTTACTGTCACACTAAACAGACGTATGTTAGC
TTCTGGATGGCTATGCTATGAATTTCATGGTCTCTAAATGCAGAGGAAGCCCCGGAAACTACC
GAAGTCACTCTCATTACCTCCCCCTCTTTCTTATATCAGAGAAGATGACT**TGAATGCATTAG**
(SEQ ID NO:32)



FIGURE 7A

Alignment Report of alignment.DNA, using Clustal method with Weighted residue weight table.
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1 ATGGC - CAAACGAGCTCGGCTAACAGCAGCTCCCTCAATCCGGTCTACCCCT Fiber 16 Genbank (SEQ ID NO:33)
1 ATGAAGC[GCC]CAAGACCGTCT[GAA-GATA]ACCTTCAA[C]CC[G]T[G]TA[T]CC[A]T Ad5/16 Fiber DNA (SEQ ID NO:38)

50 ATGAAGATGAAAGCAGCTCACAAACACCCCTTATAAAACCCGGTTTCATT Fiber 16 Genbank
50 ATGAAGATGAAAGCAGCTCACAAACACCCCTTATAAAACCCGGTTTCATT Ad5/16 Fiber DNA

100 TCCCTCAAATGGTTTGCAACAAAGCCCAGATGGAGTTCTAACCTCTTAAATG Fiber 16 Genbank
100 TCCCTCAAATGGTTTGCAACAAAGCCCAGATGGAGTTCTAACCTCTTAAATG Ad5/16 Fiber DNA

150 TGTTAATCCACTCACCTACCGCCAGCGGACCCCTCCAACTTAAAGTTGGAA Fiber 16 Genbank
150 TGTTAATCCACTCACCTACCGCCAGCGGACCCCTCCAACTTAAAGTTGGAA Ad5/16 Fiber DNA

200 GCAGTCTTACAGTAGATACTATCGATGGGTCTTGGAGGAAAATAACT Fiber 16 Genbank
200 GCAGTCTTACAGTAGATACTATCGATGGGTCTTGGAGGAAAATAACT Ad5/16 Fiber DNA

250 GCGGCAGGCCACTCACAAACTAACCAACTCCATAGGTTATTAAATAGG Fiber 16 Genbank
250 GCG[G]AGGCCACTCACAAACTAACCAACTCCATAGGTTATTAAATAGG Ad5/16 Fiber DNA

300 ATCTGGCTTGCAAAACAAAGGATGATAAAACTTGTATTATCGCTGGGAGATG Fiber 16 Genbank
300 ATCTGGCTTGCAAAACAAAGGATGATAAAACTTGTATTATCGCTGGGAGATG Ad5/16 Fiber DNA

350 GGTTGGTAAACAAAGGATGATAAAACTATGTTATCGCTGGGAGATGGGTTA Fiber 16 Genbank
350 GGTTGGTAAACAAAGGATGATAAAACTATGTTATCGCTGGGAGATGGGTTA Ad5/16 Fiber DNA

400 ATAACAAAAAATGATGTACTATGTGCCAAACTAGGACATGGCCTTGTGTT Fiber 16 Genbank
400 ATAACAAAAAATGATGTACTATGTGCCAAACTAGGACATGGCCTTGTGTT Ad5/16 Fiber DNA

450 TGACTCTCCAATGCTATCACCATAGAAAAACAACACCTTGTGGACAGGGC Fiber 16 Genbank
450 TGACTCTCCAATGCTATCACCATAGAAAAACAACACCTTGTGGACAGGGC Ad5/16 Fiber DNA

500 CAAACCAAGGCCAACTGTGTAATTAAAGAGGGAGAAGATTCAGAC Fiber 16 Genbank
500 CAAACCAAGGCCAACTGTGTAATTAAAGAGGGAGAAGATTCAGAC Ad5/16 Fiber DNA

550 TGTAAGCTCACTTAGTTCTAGTGAAGAATGGAGGACTGATAAATGGATA Fiber 16 Genbank
550 TGTAAGCTCACTTAGTTCTAGTGAAGAATGGAGGACTGATAAATGGATA Ad5/16 Fiber DNA

600 CATAACATTAATGGGAGCCTCAGAATATACTAACACCTTGTATTAAACAC Fiber 16 Genbank
600 CATAACATTAATGGGAGCCTCAGAATATACTAACACCTTGTATTAAACAC Ad5/16 Fiber DNA

650 ATCAAGTTACAATCGATGTAACCTCGCATTGATAATACTGGCCAAATT Fiber 16 Genbank
650 ATCAAGTTACAATCGATGTAACCTCGCATTGATAATACTGGCCAAATT Ad5/16 Fiber DNA

700 ATTACTTACCTATCATCCCTAAAGTAACCTGAACCTTAAAGACAACCA Fiber 16 Genbank
700 ATTACTTACCTATCATCCCTAAAGTAACCTGAACCTTAAAGACAACCA Ad5/16 Fiber DNA

750 AACATGGCTACTGGAACCCATAACCAAGTGCCTAACGGCTTCATGCCAGCA Fiber 16 Genbank
750 AACATGGCTACTGGAACCCATAACCAAGTGCCTAACGGCTTCATGCCAGCA Ad5/16 Fiber DNA

800 CCACCGCCTATCCATTATAACATACGCCACTGAGACCCCTAAATGAGAT Fiber 16 Genbank
800 CCACCGCCTATCCATTATAACATACGCCACTGAGACCCCTAAATGAGAT Ad5/16 Fiber DNA

850 TACATTATGGAGAGTGTACTACAAATCTACCAATGGAACTCTCTTCC Fiber 16 Genbank
850 TACATTATGGAGAGTGTACTACAAATCTACCAATGGAACTCTCTTCC Ad5/16 Fiber DNA

900 ACTAAAAGTTACTGTCACACTAAACAGACGTATGTTAGCTCTGGAAATGG Fiber 16 Genbank
900 ACTAAAAGTTACTGTCACACTAAACAGACGTATGTTAGCTCTGGAAATGG Ad5/16 Fiber DNA

950 CCTATGCTATGAATTTCATGGTCTCTCAAATGCAGAGGAAGGCCCGGAA Fiber 16 Genbank
950 CCTATGCTATGAATTTCATGGTCTCTCAAATGCAGAGGAAGGCCCGGAA Ad5/16 Fiber DNA

1000 ACTACCGAAGTCACTCTCATTCACCTCCCCCTCTTTTTCTTATATCAG Fiber 16 Genbank
1000 ACTACCGAAGTCACTCTCATTCACCTCCCCCTCTTTTTCTTATATCAG Ad5/16 Fiber DNA

1050 AGAAGATGACTGA
1050 AGAAGATGACTGAATGCATTAGTTGTGTTATGTTCAACGTGTTATT Fiber 16 Genbank
1050 AGAAGATGACTGAATGCATTAGTTGTGTTATGTTCAACGTGTTATT Ad5/16 Fiber DNA

1062 TCAATTG Fiber 16 Genbank
1100 TCAATTG Fiber 16 Genbank
Ad5/16 Fiber DNA

Decoration 'Decoration #1': Box residues that differ from Fiber 16 Genbank.

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FIGURE 7B

Alignment Report of alignment.prot, using Clustal method with PAM250 residue weight table.
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1	M A K R A R L S S - S F N P V Y P Y E D E S S S Q H P F I N	genbank Fiber 16 (SEQ ID NO:35)
1	M [] K R A R P S E D T F N P V Y P Y E D E S S S Q H P F I N	chimeric Ad5/Fib16 (SEQ ID NO:34)
30	P G F I S S N G F A Q S P D G V L T L K C V N P L T T A S G	genbank Fiber 16
30	P G F I S S N G F A Q S P D G V L T L K C V N P L T T A S G	chimeric Ad5/Fib16
60	P L Q L K V G S S L T V D T I D G S L E E N I T A A A P L T	genbank Fiber 16
60	P L Q L K V G S S L T V D T I D G S L E E N I T A A A P L T	chimeric Ad5/Fib16
90	K T N H S I G L L I G S G L Q T K D D K L C L S L G D G L V	genbank Fiber 16
90	K T N H S I G L L I G S G L Q T K D D K L C L S L [] E D G L V	chimeric Ad5/Fib16
120	T K D D K L C L S L G D G L I T K N D V L C A K L G H G L V	genbank Fiber 16
120	T K D D K L C L S L G D G L I T K N D V L C A K L G H G L V	chimeric Ad5/Fib16
150	F D S S N A I T I E N N T L W T G A K P S A N C V I K E G E	genbank Fiber 16
150	F D S S N A I T I E N N T L W T G A K P S A N C V I K E G E	chimeric Ad5/Fib16
180	D S P D C K L T L V L V K N G G L I N G Y I T L M G A S E Y	genbank Fiber 16
180	D S P D C K L T L V L V K N G G L I N G Y I T L M G A S E Y	chimeric Ad5/Fib16
210	T N T L F K N N Q V T I D V N L A F D N T G Q I I T Y L S S	genbank Fiber 16
210	T N T L F K N N Q V T I D V N L A F D N T G Q I I T Y L S S	chimeric Ad5/Fib16
240	L K S N L N F K D N Q N M A T G T I T S A K G F M P S T T A	genbank Fiber 16
240	L K S N L N F K D N Q N M A T G T I T S A K G F M P S T T A	chimeric Ad5/Fib16
270	Y P F I T Y A T E T L N E D Y I Y G E C Y Y K S T N G T L F	genbank Fiber 16
270	Y P F I T Y A T E T L N E D Y I Y G E C Y Y K S T N G T L F	chimeric Ad5/Fib16
300	P L K V T V T L N R R M L A S G M A Y A M N F S W S L N A E	genbank Fiber 16
300	P L K V T V T L N R R M L A S G M A Y A M N F S W S L N A E	chimeric Ad5/Fib16
330	E A P E T T E V T L I T S P F F F S Y I R E D D	genbank Fiber 16
330	E A P E T T E V T L I T S P F F F S Y I R E D D	chimeric Ad5/Fib16

Decoration 'Decoration #1': Box residues that differ from genbank Fiber 16 .